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RVPVNGENKANVASPPSIBEVVAPDSAATISISDKAPESVVPAEKPPSSGSNFVVS
ASAPRULDIDSDVEPERIKGANIVERAPNPKALSPPAAPQEDLWDFKKYIGFEEPVE
AKDDGWAVADDAGSFEHHQNHDSGPLAGENVMNVVVVAAECSPWCKTGGLGDVAGALP
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PLFRIRKOEDIYGGSROEIMKRMILECKAAVEVPWHVPCGGVPYGDGMIJVFIANDWHTA
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PDRAEAHKLIEALGHCLRTYRDYRESWRGLQERGNSQDFSWEHAAKLYEDVLLKAKYO
                                                                                                                                                                             Triticum aestivum starch synthase IIA mRNA, complete cds. API55217
AFI55217 AFI55217 AFI55217 GI:5834420
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Pooideae; Triticae; Triticum.
1 (bases 1 to 2842)
Li,Z., Chu,X., Moullle,G., Yan,L., Kosar-Hashemi,B., Hey,S.,
Napier,J., Shewry,P., Clarke,B., Appels,R., Morell,M.K. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Direct Submission
Submitted (07-SEP-1999) CSIRO Division of Plant Industry, GPO Box Submitted (07-SEP-1999) CSIRO Division of Plant Industry, GPO Box 1600, Canberra, ACT 2601, Australia Sequence update by submitter On Sep 7, 1999 this sequence version replaced gi:5825479.

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (30-MAY-1999) CSIRO Division of Plant Industry, GPO Box 1600, Canberra, ACT 2601, Australia (base 1 to 2842) Li.2., Chu.X., Moullle,G., Yan,L., Kosar-Hashemi,B., Hey,S., Napler,J., Shewry,P., Clarke,B., Appels,R., Morell,M.K. and
Li,z., Chu,X., Mouille,G., Yan,L., Kosar-Hashemi,B., Hey,S.,
Napler,J., Shewry,P., Clarke,B., Appels,R., Morell,M.K. and
Balman,S.
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                                                                                 GTCCCCCCCCCCCGCCATT 4511
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/codon_start=1
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Pooideae; Triticaee; Triticum.
I (bases 1 to 2825)
Block, M. and Loerz, H.
NUCLEIC ACID MOLECULES ENCODING ENZYMES FROM WHEAT WHICH ARE
INVOLVED IN STARKH SYNTHESIS
Patent: WO 9745545-A 5 04-DEC-1997;
HOECHST SCHERING AGREYO GMBH (DE); BLOCK MARTINA (DE)
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/clone_lib="CDNA LIBRARY IN PBLUESCRIPT SK (-)"
162. 2561
/note="unnamed protein product"
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Score 354.4; DB
Pred. No. 5e-34;
0; Mismatches
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Sequence 5 from Patent W09745545.
A93354 1 GI:6741621
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